

10/541,086

Sequence Alignment

RESULT 14

US-09-946-374-372

; Sequence 372, Application US/09946374

; Publication No. US20030073129A1

; GENERAL INFORMATION:

; APPLICANT: Baker, Kevin P.

; APPLICANT: Botstein, David

; APPLICANT: Desnoyers, Luc

; APPLICANT: Eaton, Dan L.

; APPLICANT: Ferrara, Napoleone

; APPLICANT: Fong, Sherman

; APPLICANT: Gao, Wei-Qiang

; APPLICANT: Goddard, Audrey

; APPLICANT: Godowski, Paul J.

; APPLICANT: Grimaldi, Christopher J.

; APPLICANT: Gurney, Austin L.

; APPLICANT: Hillan, Kenneth J.

; APPLICANT: Pan, James

; APPLICANT: Paoni, Nicholas F.

; APPLICANT: Roy, Margaret Ann

; APPLICANT: Smith, Victoria

; APPLICANT: Stewart, Timothy A.

; APPLICANT: Tumas, Daniel

; APPLICANT: Watanabe, Colin K.

; APPLICANT: Williams, P. Mickey

; APPLICANT: Wood, William I.

; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic

; TITLE OF INVENTION: Acids Encoding the Same

; FILE REFERENCE: P2830P1C1

Query Match 89.8%; Score 1168; DB 3; Length 269;

Best Local Similarity 94.7%; Pred. No. 1e-101;

Matches 232; Conservative 3; Mismatches 8; Indels 2; Gaps 1;

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Db      1  MAAASAGATRLLLLLLMAVAAPSRARGSGCRAGTGARGAGAEGREGEACGTVGLLLEHSF 60

Qy     61  EIDDSANFRKRGSLLNQDGTLSLSQRQLSEEEGRGLRDVAALNGLYRVIPRRPGALD 120
        |||||||
Db     61  EIDDSANFRKRGSLLNQDGTLSLSQRQLSEEEGRGLRDVAALNGLYRVIPRRPGALD 120

Qy     121 GLEAGGYVSSFPACSLVESHLSDDLTLHVDVAGNVVGVSVVTHPGGCRGHEVEDVDLEL 180
        |||||||
Db     121 GLEAGGYVSSFPACSLVESHLSDDLTLHVDVAGNVVGVSVVTHPGGCRGHEVEDVDLEL 180

Qy     181 FNTSVQLQPPPTAPGPETAFAFIERLEMEQAQKAKNPQEQKSFFAKYWHIILGGAVLLTAL 240
        |||||||
Db     181 FNTSVQLQPPPTAPGPETAFAFIERLEMEQAQKAKNPQEQKSFFAKYWHIILGGAVLLTAL 238

Qy     241 RPAAP 245
        ||
Db     239 MSGAP 243

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